AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1. (Currently Amended) A mouse comprising in its genome a first exogenous DNA molecule that functionally disrupts a NFATp gene of said mouse and a second exogenous DNA molecule that functionally disrupts a NFAT4 gene of said mouse, wherein said mouse exhibits a non-wildtype phenotype characterized by increased Th2 cytokine production relative to a wildtype mouse.

2.-32. (Canceled)

- 33. (Currently Amended) The transgenie mouse of claim <u>1</u> <u>32, 49, and 52</u>, wherein the phenotype of said mouse is <u>further</u> characterized by lymphadenopathy relative to a wild type mouse.
- 34. (Currently Amended) The transgenie mouse of claim <u>1</u> <u>32, 49, and 52</u>, wherein the phenotype of said mouse is <u>further</u> characterized by splenomegaly relative to a wild type mouse.
- 35. (Currently Amended) The transgenie mouse of claim <u>1</u>32, 49, and 52, wherein the phenotype of said mouse is <u>further</u> characterized by <u>blepharatis</u> <u>blepharitis</u> <u>relative to a wild-type mouse</u>.
- 36. (Currently Amended) The transgenie mouse of claim <u>1_32, 49, and 52</u>, wherein the phenotype of said mouse is <u>further</u> characterized by interstitial pneumonitis relative to a wild type mouse.

37. (Currently Amended) The transgenie mouse of claim <u>1</u>32, 49, and 52, wherein said mouse <u>further</u> displays an increase in peripheral T cells relative to a wild type mouse.

38. (Currently Amended) The transgenie mouse of claim 37, wherein said peripheral T cells of the mouse have a memory/activated phenotype characterized by decreased levels of Mel-14 and CD45RB, and increased levels of CD44 and CD69 relative to those of a wild-type mouse.

39. (Canceled)

- 40. (Currently Amended) The transgenie mouse of claim 139, wherein said mouse displays the peripheral T cells of the mouse display defective apoptosis relative to those of a wild-type mouse.
- 41. (Currently Amended) The transgenic mouse of claim 1, 32, 49, and 52, wherein said mouse displays the peripheral T cells of the mouse display increased Th2 cytokine production relative to those of a wild-type mouse.
- 42. (Currently Amended) The transgenic mouse of claim 41, wherein said Th2 cytokine is IL-4.

43. (Canceled)

44. (Currently Amended) The transgenie mouse of claim 43-1, wherein said mouse exhibits a higher level of antibodies having the immunoglobulin isotypes are IgG1 and IgE.

45. (Currently Amended) A method for identifying a test compound that modulates immune <u>Th2</u> cell activation via a pathway that does not involve <u>directly modulate NFATp</u> or NFAT4 comprising:

- a) administering said test compound to a first transgenic mouse comprising a genome deficient in NFATp and NFAT4;
- b) administering an appropriate control compound to a second transgenic mouse comprising a genome deficient in NFATp and NFAT4, wherein the phenotype phenotypes of the first transgenic mouse and the second transgenic mouse is are characterized by increased Th2 cytokine production[[,]]; and
- c) evaluating Th2 cell activity in said first transgenie mouse relative to Th2 cell activity in said second transgenie mouse to thereby wherein a change in Th2 cell activity in said first mouse relative to Th2 cell activity said second mouse identify identifies a compound that as one that regulates immune Th2 cell activation via a pathway that does not involve directly modulate NFATp or NFAT4.

46.-48. (Canceled)

- 49. (Currently Amended) A method for producing a transgenic mouse <u>lacking</u>

 NFATp and NFAT4, wherein said mouse exhibits a <u>non-wildtype</u> phenotype characterized by increased Th2 cytokine production relative to a corresponding wild-type mouse, comprising:
- (1<u>a</u>) introducing an exogenous DNA molecule comprising at least a portion of a NFATp gene into a mouse embryonic stem cell such that to create a first modified stem cell in which the wild type endogenous NFATp gene of the embryonic stem cell is functionally disrupted;
- (2b) introducing said transgenic mouse embryonic first modified stem cell into a pseudopregnant mouse such that said pseudopregnant mouse produces at least one offspring comprising a functionally disrupted NFATp gene;
- (3c) introducing an exogenous DNA molecule comprising at least a portion of a NFAT4 gene into a mouse embryonic stem cell to create a second modified stem cell in which such that the wild type endogenous NFAT4 gene of the embryonic stem cell is functionally disrupted;

([[4]]d) introducing said transgenic mouse embryonic second modified stem cell into a pseudopregnant mouse such that said pseudopregnant mouse produces at least one offspring comprising a functionally disrupted NFAT4 gene; and

- (5e) mating said at least one offspring with a functionally disrupted NFATp gene with said at least one offspring with a functionally disrupted NFAT4 gene and identifying subsequent offspring with both a functionally disrupted NFATp gene and a functionally disrupted NFAT4 gene to thereby produce a mouse lacking NFATp and NFAT4 which exhibits a non-wildtype phenotype characterized by increased Th2 cytokine production relative to a wild-type mouse.
- 50. (Currently Amended) A mouse transgenie An isolated cell from the mouse of claim 1 comprising a disrupted NFATp gene and a disrupted NFAT4 gene.
- 51. (Currently Amended) The mouse transgenie cell of claim 50, wherein said cell is selected from the group consisting of fertilized egg cells, embryonic stem cells and lymphoid cells.
- 52. (Currently Amended) A method for producing a transgenie mouse with a functionally disrupted NFATp gene and a functionally disrupted NFAT4 gene, wherein said mouse exhibits a <u>non-wildtype</u> phenotype characterized by increased Th2 cytokine production, comprising:

mating a transgenie mouse with having a functionally disrupted NFATp gene with a transgenie mouse with having a functionally disrupted NFAT4 gene and identifying subsequent progeny with both a functionally disrupted NFATp gene and a functionally disrupted NFAT4 gene, to thereby produce a transgenie mouse with a functionally disrupted NFATp gene and a functionally disrupted NFAT4 gene that exhibits a non-wildtype phenotype characterized by increased Th2 cytokine production.

53. (New) The mouse of claim 1, wherein the mouse is further characterized by:

- (a) blepharatis;
- (b) interstitial pneumonitis;
- (c) splenomegaly and lymphadenopathy; and
- (d) increased levels of serum IgG1 and IgE relative to a wildtype mouse.

54. (New) A mouse comprising in its genome a first exogenous DNA molecule that functionally disrupts a NFATp gene of said mouse and a second exogenous DNA molecule that functionally disrupts a NFAT4 gene of said mouse, wherein said mouse exhibits a non-wildtype phenotype characterized by increased Th2 cytokine production, blepharatis, interstitial pneumonitis splenomegaly and lymphadenopathy, and increased levels of serum IgG1 and IgE, relative to a wildtype mouse.